## **WEST Search History**

DATE: Monday, July 29, 2002

Set Name side by side	Query	Hit Count	Set Name result set
DB = USP	PT,PGPB; PLUR=YES; OP=ADJ		
L7	L6 and 14	22	L7
L6	15 and caspase and cleavage	77	L6
L5	fluorescent protein	2468	L5
L4	L1 or l2 or L3	9124	L4
L3	(((530/350)!.CCLS.))	7184	L3
L2	(((435/219)!.CCLS.))	726	L2
L1	((435/183)!.CCLS.)	1560	L1

END OF SEARCH HISTORY



**Generate Collection** 

Print

## **Search Results -** Record(s) 1 through 10 of 22 returned.

1. Document ID: US 20020098550 A1

L7: Entry 1 of 22

File: PGPB

Jul 25, 2002

PGPUB-DOCUMENT-NUMBER: 20020098550

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020098550 A1

TITLE: Death domain containing receptor 5

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw Desc Image

2. Document ID: US 20020090683 A1

L7: Entry 2 of 22

File: PGPB

Jul 11, 2002

PGPUB-DOCUMENT-NUMBER: 20020090683

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020090683 A1

TITLE: TUMOR NECROSIS FACTOR-GAMMA

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw Description

3. Document ID: US 20020081679 A1

L7: Entry 3 of 22

File: PGPB

Jun 27, 2002

PGPUB-DOCUMENT-NUMBER: 20020081679

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020081679 A1

TITLE: NARC8 programmed cell-death-associated molecules and uses

thereof

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KWIC Draw. Desc Image

4. Document ID: US 20020076799 A1

L7: Entry 4 of 22

File: PGPB

Jun 20, 2002

PGPUB-DOCUMENT-NUMBER: 20020076799

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020076799 A1

TITLE: Compositions and methods for modulating TGF-beta signaling

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw, Desc Image

5. Document ID: US 20020073441 A1

L7: Entry 5 of 22

File: PGPB

Jun 13, 2002

PGPUB-DOCUMENT-NUMBER: 20020073441

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020073441 A1

TITLE: Compositions and methods for detecting proteolytic activity

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KMC Draw Desc Image

6. Document ID: US 20020072091 A1

L7: Entry 6 of 22

File: PGPB

Jun 13, 2002

PGPUB-DOCUMENT-NUMBER: 20020072091

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020072091 A1

TITLE: Death domain containing receptor 5

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims KWIC Draw, Desc Image

7. Document ID: US 20020064828 A1

L7: Entry 7 of 22 File: PGPB

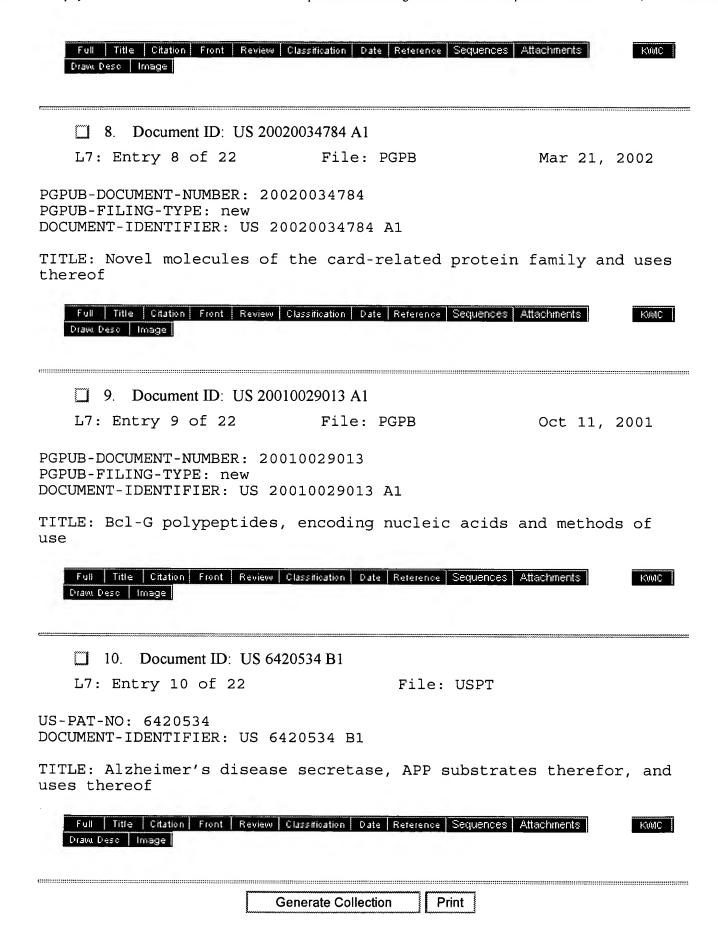
May 30, 2002

PGPUB-DOCUMENT-NUMBER: 20020064828

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020064828 A1

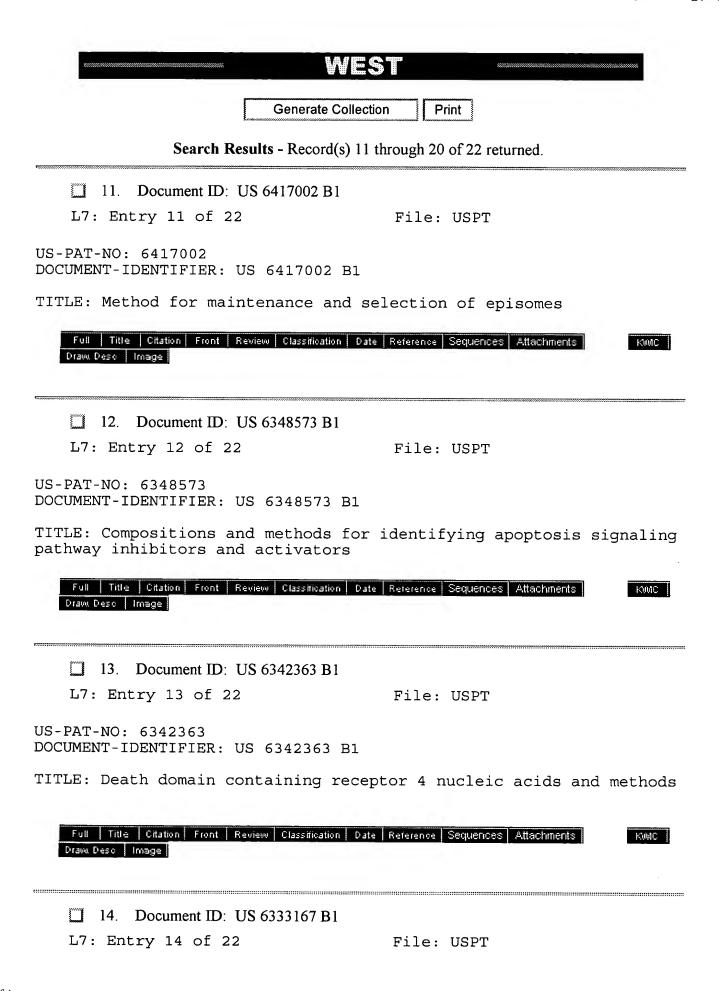
TITLE: Method of controlling the binding of calmyrin to presenilin



Terms	Documents
L6 and l4	22

Display Format: - Change Format

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US-PAT-NO: 6333167

DOCUMENT-IDENTIFIER: US 6333167 B1

TITLE: Methods and reagents for identifying inhibitors of

proteolysis of membrane-associated proteins

Full Title Citation Front Review Classification Date Reference Sequences Attachments Forward Review Draws Descriptions of the Reference Sequences Attachments

15. Document ID: US 6306663 B1

L7: Entry 15 of 22

File: USPT

US-PAT-NO: 6306663

DOCUMENT-IDENTIFIER: US 6306663 B1

TITLE: Controlling protein levels in eucaryotic organisms

Full Title Citation Front Review Classification Date Reference Sequences Attachments

Draws Description

KOMC

16. Document ID: US 6277974 B1

L7: Entry 16 of 22

File: USPT

US-PAT-NO: 6277974

DOCUMENT-IDENTIFIER: US 6277974 B1

TITLE: Compositions and methods for diagnosing and treating conditions, disorders, or diseases involving cell death



17. Document ID: US 6270964 B1

L7: Entry 17 of 22

File: USPT

US-PAT-NO: 6270964

DOCUMENT-IDENTIFIER: US 6270964 B1

TITLE: Protein fragment complementation assays for the detection of

biological or drug interactions



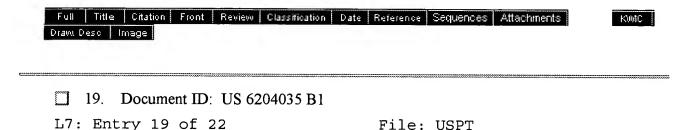
18. Document ID: US 6222017 B1

L7: Entry 18 of 22 File: USPT

US-PAT-NO: 6222017

DOCUMENT-IDENTIFIER: US 6222017 B1

TITLE: Mammalian pro-apoptotic Bok genes and their uses



File: USPT

File: USPT

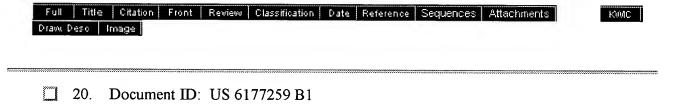
US-PAT-NO: 6204035

DOCUMENT-IDENTIFIER: US 6204035 B1

TITLE: Methods and compositions to alter the cell surface

expression of phosphatidylserine and other clot-promoting plasma

membrane phospholipids

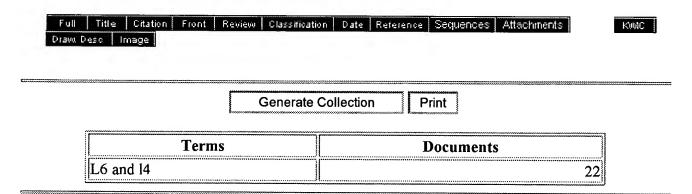


L7: Entry 20 of 22

US-PAT-NO: 6177259 DOCUMENT-IDENTIFIER: US 6177259 B1

TITLE: Assays and kits for inhibition of polyglutamine-induced cell

death



**Display Format:** -Change Format Previous Page Next Page

## **Generate Collection** Print Search Results - Record(s) 21 through 22 of 22 returned. 21. Document ID: US 6159731 A L7: Entry 21 of 22 File: USPT US-PAT-NO: 6159731 DOCUMENT-IDENTIFIER: US 6159731 A TITLE: Daxx, a Fas-binding protein that activates JNK and apoptosis Full Title Citation Front Review Classification Date Reference Sequences Attachments KWIC Draw, Desc Image 22. Document ID: US 6020143 A L7: Entry 22 of 22 File: USPT US-PAT-NO: 6020143 DOCUMENT-IDENTIFIER: US 6020143 A TITLE: Method for identifying substances that affect the interaction of a presentlin-1-interacting protein with a mammalian presenilin-1 protein Title Citation Front Review Classification Date Reference Sequences Attachments Draw, Desc - Image Print Generate Collection **Terms Documents** L6 and 14 22 Display Format: -**Change Format**

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l of l

=> d his

(FILE 'HOME' ENTERED AT 00:10:51 ON 29 JUL 2002)

	FILE 'HCAPLUS' ENTERED AT 00:10:57 ON 29 JUL 2002
L1	47626 S (PROTEIN OR POLYPEPTIDE) (L) FLUORES?
L2	1254 S L1 (L) (PROTEASE OR PROTEINASE) (L) FLUORES?
L3	45 S L2 (L) (LOOP OR SHEET)
L4	33 S L3 AND PD<19990418
L5	0 S L4 AND CASPASE
L6	41 S L2 (L) CASPASE
L7	18 S L6 AND PD<19990418

LT ANSWER 1 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1999:474278 HCAPLUS

DOCUMENT NUMBER: 131:253874

TITLE: Stoichiometry, Free Energy, and Kinetic Aspects of

Cytochrome c:Apaf-1 Binding in Apoptosis

AUTHOR(S): Purring, Cherie; Zou, Hua; Wang, Xiaodong; McLendon,

George

CORPORATE SOURCE: Department of Chemistry, Princeton University,

Princeton, NJ, 08544, USA

SOURCE: Journal of the American Chemical Society (1999

), 121(32), 7435-7436

CODEN: JACSAT; ISSN: 0002-7863

PUBLISHER: American Chemical Society

DOCUMENT TYPE: Journal LANGUAGE: English

A few years ago, a discovery was made that cytochrome c, the paradigm for a small redox protein, also plays a major role in apoptosis. It has been shown that cytochrome c (Cc) along with ATP or dATP are necessary cofactors with apoptosis **protease** activating factor-1 (Apaf-1) for the activation of caspase-9. Caspases, a family of genetically conserved cysteine proteases, are key mediators of the execution phase of apoptosis. It has been indirectly inferred that cytochrome c forms a complex with Apaf-1, but until now the direct binding of cytochrome c has not been demonstrated. We have taken the first step in characterizing this mechanism by showing that cytochrome c does, in fact, bind strongly to Apaf-1, apparently with a 2:1 Cc:Apaf-1 stoichiometry. Using fluorescence polarization methods we have measured the binding const. for Apaf-1 and horse heart ZnCc, a fluorescent deriv. of Cc in which the heme iron is replaced by zinc. This species is known to be structurally equiv. to wild-type FeCc in binding to physiol. protein partners of Cc. We conclude from all the data that Apaf-1 binds two Cc mols. with high affinity (K > 1011 M-1). The observation of such strong binding has interesting implications for the cellular control of apoptosis. If the cytosolic vol. of a typical human cell is taken to be approx. 10-11 L, then the amt. of cytochrome c that must be released from the mitochondria for Apaf-1 binding to take place involves only individual mols. If so, then Cc:Apaf-1 binding is less well described as an equil. phenomenon, but rather as a cellular signaling event. In essence every Cc mol. released from the mitochondria could bind to Apaf-1 and presumably trigger caspase-9 activation.

REFERENCE COUNT: 14 THERE ARE 14 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 2 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1999:447451 HCAPLUS

DOCUMENT NUMBER: 132:48398

TITLE: Role of acidic fibroblast growth factor in the mouse

submandibular gland carcinogenesis

AUTHOR(S): Yoshioka, Yukio

CORPORATE SOURCE: Sch. Dent., Univ. Tokushima, Tokushima, 770-8504,

Japan

SOURCE: Shikoku Shigakkai Zasshi (1999), 12(1),

133-156

CODEN: SSZAED; ISSN: 0914-6091

PUBLISHER: Shikoku Shigakkai

DOCUMENT TYPE: Journal LANGUAGE: Japanese

AB The effects of acidic fibroblast growth factor (aFGF) on the growth of mouse submandibular gland carcinoma cells and 9,10-dimethyl-1,2-benzanthracene (DMBA)-induced mouse submandibular gland carcinogenesis were examd. Recombinant human aFGF, bovine brain aFGF (bo-aFGF), recombinant human FGF-4 and 5 all stimulated the growth of mouse submandibular gland carcinoma YTcl 1-2 cells which were derived from an undifferentiated carcinoma. Goal: To det. whether aFGF could modify the

PMBA-induced mouse submandibular gland carcinogenesis. Groups 1 and 2 female mice received 1 mg of DMBA into their left submandibular glands. Thereafter, they were given a s.c. administration of either 0.5 mg/50 .mu.L bo-aFGP or vehicle, resp., three times a week for 10 wk. Groups 3 and 4 mice were given either bo-aFGF or vehicle only. Sixteen weeks after the beginning of the expt. the incidence of carcinomas in Group 1 was significantly (p <0.05) greater than that in Group 2. The mean wt. of submandibular glands in Group 1 mice was the greatest indicating that bo-aFGF can promote the development of submandibular gland carcinoma. AFGF was demonstrated in granular convoluted tubule cells and ductal cells in the normal submandibular glands and carcinomas by immunohistochem. staining. Immunoblot anal. detected protein with mol. wt. of 18 kDa in normal and tumor tissues and YTcl 1-2 cells. Expression of aFGF mRNA was also demonstrated in all exptl. samples and YTcl 1-2 cells. Immunohistochem. study and immunoblot anal. revealed the expression of FGF receptor (FGFR)-1 and -4, but not FGFR-2 and -3 in normal submandibular gland and tumor tissues, indicating that aFGF binds to these receptors of the submandibular gland carcinomas and may stimulate cell growth by autocrine and paracrine mechanisms. The growth signal of aFGF should be transmitted into cells by tyrosine phosphorylation of its receptor. Tyrphostins are artificial and low mol. wt. compds. that specifically inhibit protein tyrosine kinases. Tyrphostins 9, 23, 47, B42, and B46 suppressed the growth of YTcl 1-2 cells, and tyrphostin 9 was found to he the most potent inhibitor; the IC50 was estd. to be 0.1 .mu.M. Furthermore, the growth of nude mouse tumors induced by YTcl 1-2 cell inoculation was suppressed by intra-tumor administration of tyrphostin 9 (P < 0.001), indicating that it was effective in vitro and in vivo Tyrphostin 9 inhibited tyrosine-phosphorylation of membrane proteins including FGFR-1 in the aFGF-treated YTcl 1-2 cells Flow cytometric anal. of the tyrphostin 9-treated cell DNA indicated that tyrphostin 9 induced the arrest of cell cycle and apoptosis. Indeed, DNA fragmentation of the treated YTcl 1-2 cells was demonstrated by fluorescent DNA staining with Hochest 33258 and terminal deoxynucleotidyl transferase mediated dUTP-biotin nick end labeling method. Electron microscopic study also revealed nuclear disorganization, development of cytoplasmic vacuoles, and decrease of mitochondria in the tyrphostin-treated cells. The dysfunction of mitochondria membrane was also indicated by the leakage of incorporated Rhodamin 123 from mitochondria. Dose- and time-dependent activation of apoptosis-assocd. protease, caspase-3, was detected after the treatment with this tyrphostin. These results suggest that aFGF plays a role in the development of mouse submandibular gland carcinomas and the growth of carcinoma cells, and that tyrosine kinase inhibitor, tyrphostin 9, can be used as an inhibitor for the submandibular gland carcinoma.

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ANSWER 3 OF 18 HCAPLUS COPYRIGHT 2002 ACS
ACCESSION NUMBER:
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DOCUMENT NUMBER:

1999:397367 HCAPLUS 131:181483

TITLE:

Novel mutant green fluorescent protein protease substrates reveal

the activation of specific caspases during

apoptosis

AUTHOR(S):

Mahajan, Nupam P.; Harrison-Shostak, D. Corinne;

Michaux, Jennifer; Herman, Brian

CORPORATE SOURCE:

The Lineberger Comprehensive Cancer Center and

Department of Cell Biology and Anatomy, University of North Carolina at Chapel Hill, Chapel Hill, NC, 27599,

SOURCE:

PUBLISHER:

Chemistry & Biology (1999), 6(6), 401-409

CODEN: CBOLE2; ISSN: 1074-5521 Current Biology Publications

DOCUMENT TYPE:

Journal LANGUAGE: English

The caspase-mediated proteolysis of many cellular proteins is a crit. event during programmed cell death or apoptosis. It is important to det. which caspases are activated in mammalian cells, and where and when activation occurs, upon receipt of specific death stimuli.

information would be useful in the design of strategies to regulate the activation of caspases during apoptosis. We developed two novel fluorescent substrates that were specifically cleaved by caspase-1 or caspase-3. For in vitro studies, four-amino-acid recognition sequences, YVAD for caspase-1 and DEVD for caspase-3, were introduced between blue fluorescent protein (BFP) and green fluorescent protein (GFP), expressed in bacteria and purified. For in vivo studies, YVAD and DEVD were introduced between cyan fluorescent protein and yellow fluorescent protein, and expression was monitored in live mammalian cells. proximity between fluorophores was detd. using fluorescence resonance energy transfer. Purified substrates were cleaved following exposure to purified caspase-1 and caspase-3. In Cos-7 cells, caspase-1 and caspase-3 substrates were cleaved upon induction of apoptosis with staurosporine, a protein-kinase inhibitor, whereas caspase-3 but not caspase-1 substrate was cleaved upon treatment of cells with the DNA-damaging agent mitomycin c. These substrates allow the spatial activation of specific members of the caspase family to be deciphered during the initiation and execution phase of programmed cell death, and allow activation of specific caspases to be monitored both in vivo and in vitro. This technol. is also likely to be useful for high-throughput screening of reagents that modulate caspase activity.

REFERENCE COUNT: 18 THERE ARE 18 CITED REFERENCES AVAILABLE FOR THIS

RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 4 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1999:378030 HCAPLUS

DOCUMENT NUMBER: 131:179425

TITLE: Inhibition of caspase activity does not prevent the

signaling phase of apoptosis in prostate cancer cells

AUTHOR(S): Denmeade, Samuel R.; Lin, Xiaohui S.; Tombal,

Bertrand; Isaacs, John T.

CORPORATE SOURCE: Johns Hopkins Oncology Center, Johns Hopkins

University School of Medicine, Baltimore, MD, USA

Prostate (New York) (1999), 39(4), 269-279

CODEN: PRSTDS; ISSN: 0270-4137

PUBLISHER: Wiley-Liss, Inc.

DOCUMENT TYPE: Journal LANGUAGE: English

SOURCE:

BACKGROUND. Caspases are a family of cysteine proteases capable of characteristically cleaving after an aspartic acid residue. Various members of the caspase family (e.g., caspases 8 and 9) have been implicated as crit. initiators in the signaling phase, while others (e.g., caspases 3, 6, and 7) have been implicated in the effector or execution phase of apoptosis. Thapsigargin (TG) is capable of inducing cell proliferation-independent apoptosis of prostate cancer cells. This study was undertaken to det. if caspase inhibition can prevent TG- or 5-fluorodeoxyuridine (5-FrdU)-induced apoptosis in prostate cancer cells. METHODS. Caspase activity was evaluated by Western blot anal. of the cleavage of retinoblastoma (Rb) protein, a caspase substrate during TG-induced death of prostate cancer cells. In addn., hydrolysis of caspase-specific fluorescent peptide substrates was assayed in lysates from TG-treated cells. Clonogenic survival assays were performed following treatment of rat AT3 and human TSU-Pr1 prostate cancer cell lines with TG and 5-FrdU in the presence and absence of peptide caspase inhibitors. AT3.1 cells transfected with the crmA gene, encoding a viral protein with caspase-inhibitory activity, were also tested for clonogenic survival following TG and 5-FrdU exposure. During treatment with TG, Rb is first dephosphorylated and then proteolytically cleaved into 100-kDa and 40-kDa forms, indicative of caspase activity. A 6-8-fold increase in class II (i.e., caspases 3, 7, and 10) hydrolysis of the caspase substrate Z-DEVD-AFC was obsd. after 24 h of TG or 5-FrdU. AT3 cells expressing crmA (i.e., an inhibitor of caspases 1, 4, and 8) were not protected from apoptosis induced by TG or 5-FrdU. caspase inhibitors Z-DEVD-fmk (i.e., an inhibitor of caspases 3, 7, and 10) and Z-VAD-fmk (i.e., a general

caspase inhibitor) were also unable to protect TSU and AT3 cells from apoptosis induced by TG or 5-FrdU. CONCLUSIONS. Caspase activation may play a role in the downstream effector phase of the apoptotic cascade; however, in this study, caspase inhibition did not prevent the signaling phase of apoptosis induced by two agents with distinct mechanisms of cytotoxicity, TG or 5-FrdU. These results suggest that caspase inhibition by recently described endogenous caspase inhibitors should not lead to development of resistance to TG. A strategy for targeting TG's unique cytotoxicity to metastatic prostate cancer cells is currently under development.

REFERENCE COUNT: 58 THERE ARE 58 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 5 OF 18 HCAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1999:336264 HCAPLUS

DOCUMENT NUMBER: 131:100472

TITLE: Localization, regulation and possible consequences of

> apoptotic protease-activating factor-1 (Apaf-1) expression in granulosa cells of the mouse ovary

Robles, Rodolfo; Tao, Xiao-Jing; Trbovich, Alexander AUTHOR(S):

M.; Maravei, Daniel V.; Nahum, Ravit; Perez, Gloria

I.; Tilly, Kim I.; Tilly, Jonathan L.

Vincent Center for Reproductive Biology, Department of CORPORATE SOURCE:

Obstetrics and Gynecology, Massachusetts General

Hospital/Harvard Medical School, Boston, MA,

02114-2696, USA

Endocrinology (1999), 140(6), 2641-2644 CODEN: ENDOAO; ISSN: 0013-7227 SOURCE:

PUBLISHER: Endocrine Society

DOCUMENT TYPE: Journal LANGUAGE: English

The recent characterization of apoptotic protease-activating factor-1 (Apaf-1) in vertebrates as a putative homolog of the Caenorhabditis elegans gene, ced-4, indicates that the third major arm of the C. elegans programmed cell death machinery has also been conserved through evolution. Although apoptosis is now known to be important for ovarian follicular atresia in vertebrates, nothing is known of the role of Apaf-1 in ovarian function. Herein the authors show by immunohistochem. anal. that Apaf-1 is abundant in granulosa cells of early antral follicles, whereas in vivo gonadotropin priming completely suppresses Apaf-1 expression and granulosa cell apoptosis. Western blot anal. of fractionated protein exts. prepd. from granulosa cells before and after in vitro culture without hormonal support to induce apoptosis indicated that mitochondrial cytochrome c release, a biochem. step required for the activation of Apaf-1, occurs in granulosa cells cultured in vitro. Moreover, Western blot anal. of procaspase-3 processing, a principal downstream event set in motion by activated Apaf-1, indicated that healthy granulosa cells possess almost exclusively the inactive (pro-) form of the enzyme, whereas granulosa cells deprived of hormonal support rapidly process procaspase-3 to the active enzyme. Lastly, the authors show that serum-starved granulosa cells activate caspase -3-like enzymes both prior to and after nuclear pyknosis, as revealed by a single-cell fluorescent caspase activity assay. These data, combined with previous observations regarding the role of homologs of the two other C. elegans cell death regulatory genes, ced-9 (Bcl-2 family members) and ced-3 (caspases), in atresia fully support the hypothesis that granulosa cell apoptosis is precisely coordinated by all three major arms of a cell death program conserved through evolution. THERE ARE 18 CITED REFERENCES AVAILABLE FOR THIS REFERENCE COUNT: 18 RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 6 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1999:309578 HCAPLUS

DOCUMENT NUMBER: 131:100642

TITLE: Caspase-mediated cleavage of cytoskeletal actin plays

a positive role in the process of morphological

apoptosis

AUTHOR(S): Mashima, Tetsuo; Naito, Mikihiko; Tsuruo, Takashi CORPORATE SOURCE: Laboratory of Biomedical Research, Institute of

Molecular and Cellular Biosciences, University of

Tokyo, Tokyo, 113, Japan

SOURCE: Oncogene (1999), 18(15), 2423-2430

CODEN: ONCNES; ISSN: 0950-9232

PUBLISHER: Stockton Press

DOCUMENT TYPE: Journal LANGUAGE: English

Tumors result from the imbalance between cell growth and apoptosis. of the characteristic changes in cancers is the abnormality in cytoskeleton, which suggests some roles of cytoskeletal proteins in tumorigenesis or the maintenance of tumor cells. Previously the authors showed that cytoskeletal actin is the substrate of caspases, the proteases responsible for apoptosis, while the role of actin cleavage in apoptosis remained unknown. To examine the cleavage of actin in vivo, the authors extensively performed immunoblot anal. using actin fragment-specific antibody. Here, the authors showed that, in some solid tumor cells, induction of apoptosis was accompanied by caspase-dependent actin-cleavage to 15 and 31 kDa fragments in vivo. To elucidate the role of actin-cleavage further, the authors introduced actin cleaved-fragments. The authors found that ectopic expression of an actin 15 kDa fragment induces morphol. changes resembling those of apoptotic cells. The expression of the actin fragment induced a dramatic change of cellular actin localization, as visualized by enhanced green fluorescent protein (EGFP)-tagged actin, while the actin fragment expression did not cause caspase activation nor the cleavage of a marker substrate protein, poly (ADP-ribose) polymerase. These results indicate that actin cleavage could play a pos. role in the morphol. changes of apoptosis downstream of caspase activation.

REFERENCE COUNT:

THERE ARE 46 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 7 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1999:242174 HCAPLUS

DOCUMENT NUMBER: 131:72217

TITLE: Expression of extended polyglutamine sequentially

activates initiator and effector caspases

AUTHOR(S): Miyashita, Toshiyuki; Matsui, Jun; Ohtsuka, Yuko; U,

Mami; Fujishima, Sayaka; Okamura-Oho, Yuko; Inoue,

Tadashi; Yamada, Masao

CORPORATE SOURCE: Department of Genetics, National Children's Medical

Research Center, Tokyo, 154-8509, Japan

SOURCE: Biochemical and Biophysical Research Communications (

**1999**), 257(3), 724-730

CODEN: BBRCA9; ISSN: 0006-291X

PUBLISHER: Academic Press

DOCUMENT TYPE: Journal LANGUAGE: English

To date, eight neurodegenerative disorders, including Huntington's disease and dentatorubral-pallidoluysian atrophy, have been identified to be caused by expansion of a CAG repeat coding for a polyglutamine (polyQ) stretch. It is, however, unclear how polyQ expansion mediates neuronal cell death obsd. in these disorders. Here, the authors have established a tetracycline-regulated expression system producing 19 and 56 repeats of glutamine fused with green fluorescent protein.

Induced expression of the 56 polyQ stretch, but not of the 19 polyQ

Induced expression of the 56 polyQ stretch, but not of the 19 polyQ stretch, caused marked nuclear aggregation and apoptotic morphol. changes of the nucleus. In vitro enzyme assays and Western blotting showed that polyQ56 expression sequentially activated initiator and effector

caspases, such as caspase-8 or -9, and caspase

-3, resp. Furthermore, using cell-permeable fluorogenic substrate, the activation of caspase-3-like proteases was demonstrated in intact cells with aggregated polyQ. This is the first

direct evidence that the expression of extended polyQ activates caspases and together with the previous findings that some of the

products of genes responsible for CAG repeat diseases are substrates of caspase-3 indicates an important role of caspases in the

pathogenesis of these diseases. (c) 1999 Academic Press.

REFERENCE COUNT: 28 THERE ARE 28 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 8 OF 18 HCAPLUS COPYRIGHT 2002 ACS T.7 ACCESSION NUMBER: 1999:37963 HCAPLUS

DOCUMENT NUMBER: 130:164184

Toxic bile salts induce rodent hepatocyte apoptosis TITLE:

via direct activation of Fas

AUTHOR(S): Faubion, William A.; Guicciardi, M. Eugenia; Miyoshi,

Hideyuki; Bronk, Steven F.; Roberts, Patricia J.; Svingen, Phyllis A.; Kaufmann, Scott H.; Gores,

Gregory J.

CORPORATE SOURCE: Division of Gastroenterology and Hepatology, Mayo

Medical School, Clinic, and Foundation, Rochester, MN,

55905, USA

SOURCE: Journal of Clinical Investigation (1999),

103(1), 137-145

CODEN: JCINAO; ISSN: 0021-9738

DOCUMENT TYPE:

PUBLISHER: American Society for Clinical Investigation

LANGUAGE: English

Cholestatic liver injury appears to result from the induction of hepatocyte apoptosis by toxic bile salts such as glycochenodeoxycholate (GCDC). Previous studies from this lab. indicate that cathepsin B is a downstream effector protease during the hepatocyte apoptotic process. Because caspases can initiate apoptosis, the present studies were undertaken to det. the role of caspases in cathepsin B activation. Immunoblotting of GCDC-treated McNtcp.24 hepatoma cells demonstrated cleavage of poly(ADP-ribose) polymerase and lamin B1 to fragments that indicate activation of effector caspases. Transfection with CrmA, an inhibitor of caspase 8, prevented GCDC-induced cathepsin B activation and apoptosis. Consistent with these results, an increase in caspase 8-like activity was obsd. in GCDC-treated cells. Examn. of the mechanism of GCDC-induced caspase 8 activation revealed that dominant-neg. FADD inhibited apoptosis and that hepatocytes isolated from Fas-deficient lymphoproliferative mice were resistant to GCDC-induced apoptosis. After GCDC treatment, immunopptn. expts. demonstrated Fas oligomerization, and confocal microscopy demonstrated .DELTA.FADD-GFP (Fas-assocd. death domain-green fluorescent protein), aggregation in the absence of detectable Fas ligand mRNA. Collectively, these data suggest that GCDC-induced hepatocyte apoptosis involves ligand-independent oligomerization of Fas, recruitment of FADD, activation of caspase

downstream caspases and cathepsin B. REFERENCE COUNT: 42 THERE ARE 42 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

8, and subsequent activation of effector proteases, including

ANSWER 9 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1998:797528 HCAPLUS

DOCUMENT NUMBER: 130:135767

AUTHOR(S):

SOURCE:

TITLE: TRAIL/Apo2L activates c-Jun NH2-terminal kinase (JNK)

via caspase-dependent and caspase-independent pathways Muhlenbeck, Frank; Haas, Elvira; Schwenzer, Ralph;

Schubert, Gisela; Grell, Matthias; Smith, Craig;

Scheurich, Peter; Wajant, Harald

CORPORATE SOURCE: Institute of Cell Biology and Immunology, University

of Stuttgart, Stuttgart, 70569, Germany Journal of Biological Chemistry (1998),

273(49), 33091-33098

CODEN: JBCHA3; ISSN: 0021-9258

PUBLISHER: American Society for Biochemistry and Molecular

Biology

DOCUMENT TYPE: Journal LANGUAGE: English

In this study, we show that TRAIL (tumor necrosis factor-related apoptosis-inducing ligand), also called Apo2L, activates the c-Jun N-terminal kinase (JNK). Interestingly, TRAIL-induced JNK activation occurs in a cell type-specific manner. In HeLa cells, TRAIL-induced JNK activation can be completely blocked with the cysteine protease inhibitor zVAD-fmk, whereas the same inhibitor has no, or even a stimulatory, effect on JNK activation in Kym-1 cells. Hence, TRAIL can engage at least two independent pathways leading to JNK activation, one that is cysteine protease-dependent and one that is cysteine protease-independent. To investigate whether the cysteine protease-dependent signaling of TRAIL leading to JNK activation is related to the apoptotic pathway engaged by this ligand, we investigated HeLa cells stably overexpressing a dominant neg. mutant of FADD (Fas-assocg. protein with death domain) (GFP(green fluorescent protein).DELTA.FADD). In these cells, TRAIL-induced cell death and activation of the apoptosis executioner caspase-8 (FLICE/MACH) and caspase-3 (YAMA, CPP-32, Apopain), that belong to caspase subfamily of cysteine proteases, were abrogated, whereas JNK activation remained unaffected and was still sensitive toward z-VAD-fmk. Similar data were found in HeLa cells overexpressing Apol/Fas and GFP.DELTA.FADD upon stimulation with agonistic antibodies. These data suggest that crosslinking of the TRAIL receptors and Apol/Fas, resp., engages a FADD-dependent pathway leading to the activation of apoptotic caspases and, in parallel, a FADD-independent pathway leading to the stimulation of one or more cysteine proteases capable to activate JNK but not sufficient for the induction of cell death. REFERENCE COUNT: 69 THERE ARE 69 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 10 OF 18 HCAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER:

1998:766905 HCAPLUS

DOCUMENT NUMBER:

130:137744

TITLE:

Hypoxia induces apoptosis in human neuroblastoma SK-N-MC cells by caspase activation accompanying

cytochrome c release from mitochondria

AUTHOR(S):

Araya, Runa; Uehara, Takashi; Nomura, Yasuyuki

CORPORATE SOURCE: Graduate School of Pharmaceutical Sciences, Department

of Pharmacology, Hokkaido University, Sapporo,

060-0812, Japan

SOURCE:

FEBS Letters (1998), 439(1,2), 168-172

CODEN: FEBLAL; ISSN: 0014-5793

Elsevier Science B.V.

DOCUMENT TYPE:

PUBLISHER:

Journal LANGUAGE: English

We have attempted to elucidate the mechanism of apoptotic cell death induced by hypoxia (very low oxygen conditions) in neuronal cells. neuroblastoma SK-N-MC cells under hypoxic conditions resulted in apoptosis in a time-dependent manner estd. by DNA fragmentation assay and nuclear morphol. stained with fluorescent chromatin dye. Pretreatment with Z-Asp-CH2-DCB, a caspase inhibitor, suppressed the DNA ladder in response to hypoxia in a concn.-dependent manner. An increase in caspase-3-like protease (DEVDase) activity was obsd. during apoptosis, but no caspase-1 activity (YVADase) was detected. To confirm the involvement of caspase-3 during apoptosis, Western blot anal. was performed using anti-caspase-3 antibody. The 20- and 17-kDa proteins, corresponding to the active products of caspase-3, were generated in hypoxia-challenged lysates in which processing of the full length form of caspase-3 was evident. With a time course similar to this caspase-3 activation, hypoxic stress caused the cleavage of PARP, yielding an 85-kDa fragment typical of caspase activity. In addn., caspase-2 was also activated by hypoxia, and the stress elicited the release of cytochrome c into the cytosol during apoptosis. These results suggest that caspase activation and cytochrome c release play roles in hypoxia-induced neuronal apoptosis.

REFERENCE COUNT: 34 THERE ARE 34 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 11 OF 18 HCAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1998:716682 HCAPLUS

DOCUMENT NUMBER: 130:104925

TITLE: Role of Caspases in Immunotoxin-Induced Apoptosis of

Cancer Cells

AUTHOR(S): Keppler-Hafkemeyer, Andrea; Brinkmann, Ulrich; Pastan,

Ira

CORPORATE SOURCE: Laboratory of Molecular Biology Division of Basic

Sciences National Cancer Institute, National Institutes of Health, Bethesda, MD, 20892, USA

Biochemistry (1998), 37(48), 16934-16942

CODEN: BICHAW; ISSN: 0006-2960

PUBLISHER: American Chemical Society

DOCUMENT TYPE: Journal LANGUAGE: English

SOURCE:

Immunotoxins composed of antibodies linked to plant or bacterial toxins are being evaluated in the treatment of cancer. It is known that the toxin moieties of immunotoxins, including Pseudomonas exotoxin A (PE), diphtheria toxin, and ricin, are capable of inducing apoptosis. Since the efficiency of induction of apoptosis and the apoptosis pathway may have direct effects on the therapeutic usefulness of immunotoxins, we have studied how B3(Fv)-PE38, a genetically engineered immunotoxin in which the Fv fragment of an antibody is fused to a mutated form of PE, induces apoptosis of the MCF-7 breast cancer cell line. We show for the first time that a PE-contg. immunotoxin activates ICE/ced-3 proteases, now termed caspases, and causes characteristic cleavage of the "death substrate" poly(ADP)-ribose polymerase (PARP) to an 89 kDa fragment with a time course of cleavage comparable to that induced by TNF.alpha.. Also the fluorescent substrate, DEVD-AFC, is cleaved 2-4-fold more rapidly by lysates from B3(Fv)-PE38 treated MCF-7 cells than untreated control cells, suggesting that a CPP32-like caspase is involved in B3(Fv)-PE38-mediated apoptosis. B3(Fv)-PE38-induced PARP cleavage is inhibited by several protease inhibitors known to inhibit caspases (zVAD-fmk, zDEVD-fmk, zIETD-fmk) as well as by overexpression of Bcl-2 providing addnl. evidence for caspase involvement. ZVAD-fmk, a broad spectrum inhibitor of most mammalian caspases, prevents the early morphol. changes and loss of cell membrane integrity produced by B3(Fv)-PE38, but not its ability to inhibit protein synthesis, arrest cell growth, and subsequently kill cells. Despite inhibition of apoptosis, the immunotoxin is still capable of selective cell killing, which indicates that B3(Fv)-PE38 kills cells by two mechanisms: one requires caspase activation, and the other is due to the arrest of protein synthesis caused by inactivation of elongation factor 2. The fact that an immunotoxin can specifically kill tumor cells without the need of inducing apoptosis makes such agents esp. valuable for the treatment of cancers that are protected against apoptosis, e.g., by overexpression of Bcl-2.

REFERENCE COUNT: 51 THERE ARE 51 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L7 ANSWER 12 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1998:638936 HCAPLUS

DOCUMENT NUMBER: 129:341895

CORPORATE SOURCE:

TITLE: Prodomain-dependent nuclear localization of the

caspase-2 (Nedd2) precursor. A novel function for a

caspase prodomain

AUTHOR(S): Colussi, Paul A.; Harvey, Natasha L.; Kumar, Sharad

Hanson Centre for Cancer Research, Institute of

Medical and Veterinary Science, Adelaide, 5000,

Australia

SOURCE: Journal of Biological Chemistry (1998),

273(38), 24535-24542

CODEN: JBCHA3; ISSN: 0021-9258

PUBLISHER: American Society for Biochemistry and Molecular

DOCUMENT TYPE: Journal LANGUAGE: English

Caspases are cysteine proteases that play an essential role in apoptosis by cleaving several key cellular proteins. Despite their function in apoptosis, little is known about where in the cell they are localized and whether they are translocated to specific cellular compartments upon activation. In the present paper, using Aequorea victoria green fluorescent protein (GFP) fusion constructs, the authors detd. the localization of Nedd2 (mouse caspase-2) and showed that both precursor and processed caspase-2 localize to the cytoplasmic and the nuclear compartments of transfected NIH-3T3 cells. The authors demonstrated that the nuclear localization of caspase-2 was strictly dependent on the presence of the pro-domain. A caspase-2 pro-domain-GFP localized to dotand fiber-like structures, mostly in the nucleus, whereas a protein lacking the pro-domain was largely concd. in the cytoplasm. It was also shown that an N-terminal fusion of the pro-domain of caspase-2 to caspase-3 mediated nuclear transport of caspase-3, which is normally localized in the cytoplasm. These results suggest that, in addn. to roles in dimerization and recruitment through adaptors, the caspase-2 pro-domain has a novel function in nuclear transport.

L7 ANSWER 13 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1998:541778 HCAPLUS

DOCUMENT NUMBER: 129:243599

TITLE: Intracellular aggregate formation of

dentatorubral-pallidoluysian atrophy (DRPLA) protein

with the extended polyglutamine

AUTHOR(S): Miyashita, Toshiyuki; Nagao, Kazuaki; Ohmi, Kazuhiro;

Yanagisawa, Hiroko; Okamura-Oho, Yuko; Yamada, Masao

CORPORATE SOURCE: Department of Genetics and Pathology, National

Children's Medical Research Center, Tokyo, 154-8509,

Japan

SOURCE: Biochemical and Biophysical Research Communications (

**1998**), 249(1), 96-102

CODEN: BBRCA9; ISSN: 0006-291X

PUBLISHER: Academic Press

DOCUMENT TYPE: Journal LANGUAGE: English

Dentatorubral-pallidoluysian atrophy (DRPLA) is an autosomal dominant neurodegenerative disorder caused by the abnormal CAG triplet-repeat expansion resulting in an elongated polyglutamine (polyQ) stretch. have recently showed that the DRPLA protein is cleaved during apoptosis by caspase-3, one of the cysteine protease family members known to be activated during apoptosis. We report here the subcellular localization of the DRPLA protein by fusing the green fluorescent protein as a tag. The full length DRPLA protein is localized predominantly but not exclusively in the nucleus regardless of the length of the polyQ stretch. In contrast, an N-terminal-deleted fragment contg. polyQ produced by the proteolytic cleavage with caspase-3 is found both in the nucleus and the cytoplasm. Moreover, the same fragment with the elongated polyQ showed aggregation when overexpressed. Some cells with aggregate formation showed apoptotic phenotype. These findings raise the possibility that the DRPLA protein processed by caspase-3 may lead to aggregation of the protein resulting in the development of neurodegeneration. (c) 1998 Academic Press.

L7 ANSWER 14 OF 18 HCAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER: 1998:412009 HCAPLUS

DOCUMENT NUMBER: 129:202056

TITLE: Perforin-dependent nuclear entry of granzyme B

precedes apoptosis, and is not a consequence of

nuclear membrane dysfunction

AUTHOR(S): Trapani, Joseph A.; Jans, Patricia; Smyth, Mark J.;

Froelich, Christopher J.; Williams, Elizabeth A.;

Sutton, Vivien R.; Jans, David A.

CORPORATE SOURCE: Cellular Cytotoxicity Laboratory, The Austin Research

Institute, Heidelberg, 3084, Australia

Cell Death and Differentiation (1998), 5(6),

488-496

CODEN: CDDIEK; ISSN: 1350-9047

PUBLISHER: Stockton Press

DOCUMENT TYPE: Journal LANGUAGE: English

SOURCE:

Killer lymphocytes utilize the synergy of a membranolytic protein , perforin, and the serine protease granzyme B (grB) to induce target cell apoptosis, however the mechanism of this synergy remains incompletely defined. The authors have previously shown that perforin specifically induces the redistribution of cytoplasmic grB into the nucleus of dying cells, however a causal role for nuclear targeting of grB in cell death has not been demonstrated. In the present study, the authors used confocal laser scanning microscopy (CLSM) to det. whether the nuclear accumulation of fluoresceinated (FITC-) grB precedes or is a consequence of apoptosis. Two distinct and mutually exclusive cellular responses were obsd. in FDC-P1 cells: (i) up to 50% of the cells rapidly accumulated FITC-grB in the nucleus (maximal at 7 min; t1/2 of 2 min) and underwent apoptosis; (ii) the remaining cells took up FITC-grB only into the cytoplasm, and escaped apoptosis. Under these conditions, DNA fragmentation was not obsd. for at least 13 min, indicating nuclear accumulation of grB preceded the execution phase of apoptosis. Furthermore, nuclear import of grB proceeded through an intact nuclear membrane, as the nuclei of cells whose cytoplasm was pre-loaded with 70 kDa FITC-dextran excluded dextran for up to 90 min while still undergoing apoptosis in response to perforin and grB. These findings indicated that perforin-induced nuclear accumulation of grB precedes apoptosis, and is not a byproduct of caspase-induced nuclear membrane degrdn. The cell membrane lesions formed by perforin in these expts. were not large enough to permit a 13 kDa protein (yeast cdk p13suc) access into the cytoplasm, but an 8 kDa protein (bacterial azurin) was able to equilibrate between the cytosol and the exterior. Therefore, transmembrane pores large enough to allow passive diffusion of grB (32 kDa) into the cell are not necessary for apoptosis. Rather, a perforin-dependent signal results in a redistribution of grB from the cytoplasm to the nucleus, where it may contribute to the nuclear changes assocd. with apoptosis.

L7 ANSWER 15 OF 18 HCAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1998:300008 HCAPLUS

DOCUMENT NUMBER: 129:90001

TITLE: Inhibition of epidermal growth factor receptor kinase

induces protease-dependent apoptosis in human colon

cancer cells

AUTHOR(S): Karnes, William E., Jr.; Weller, Shaun G.; Adjei,

Philip N.; Kottke, Timothy J.; Glenn, Kahlil S.;

Gores, Gregory J.; Kaufmann, Scott H.

CORPORATE SOURCE: Division of Gastroenterology, Mayo Clinic, Rochester,

MN, USA

SOURCE: Gastroenterology (1998), 114(5), 930-939

CODEN: GASTAB; ISSN: 0016-5085

PUBLISHER: W. B. Saunders Co.

DOCUMENT TYPE: Journal LANGUAGE: English

AB The epidermal growth factor receptor (EGFR) is under investigation as a therapeutic target for cancers. Colon cancer cell lines are variably dependent on autocrine stimulation of EGFR. We therefore examd. the effects of a selective EGFR tyrosine kinase inhibitor, PD 153035, on proliferation and survival of five colon cancer cell lines whose autonomous proliferation is either EGFR ligand dependent or EGFR ligand independent. Effects of inhibitors were screened by MTS growth assays, [3H]thymidine incorporation, terminal deoxynucleotidyl transferase-medaited deoxyuridine triphosphate nick-end labeling assay,

fluorescence microscopy, immunoblotting, and in vitro protease assays. PD 153035 caused dose-dependent cytostasis (200 nmol/L to 1 .mu.mol/L) and apoptosis (>10 .mu.mol/L) in ligand-dependent cell lines and caused variable apoptosis (>10 .mu.mol/L) but no cytostasis in ligand-independent cell lines. Apoptosis induced by 10 .mu.mol/L PD 153035 was not assocd. with induction of p53 protein expression but was accompanied by activation of caspases that cleave poly(ADP-ribose) polymerase, lamin B1, and Bcl-2. Inhibition of caspase 3-like protease activity by DEVDfluoromethylketone significantly delayed the onset of PD 153035-induced apoptosis. The EGFR tyrosine kinase inhibitor PD 153035 induces cytostasis and caspase-dependent apoptosis in EGFR ligand-dependent colon cancer cell lines. These observations encourage further investigation of EGFR tyrosine kinase inhibitors for treatment of colorectal neoplasms.

L7 ANSWER 16 OF 18 HCAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1998:272306 HCAPLUS

DOCUMENT NUMBER:

129:25240

TITLE:

Detection of programmed cell death using fluorescence

energy transfer

AUTHOR(S):

SOURCE:

Xu, Xiang; Gerard, Amy L. V.; Huang, Betty C. B.;

Anderson, David C.; Payan, Donald G.; Luo, Ying

CORPORATE SOURCE:

Rigel, Inc., Sunnyvale, CA, 94086, USA Nucleic Acids Research (1998), 26(8),

2034-2035

CODEN: NARHAD; ISSN: 0305-1048

PUBLISHER:

Oxford University Press

DOCUMENT TYPE:

Journal

LANGUAGE:

English

Fluorescence energy transfer (FRET) can be generated when green fluorescent protein (GFP) and blue fluorescent protein (BFP) are covalently linked together by a short peptide. Cleavage of this linkage by protease completely eliminates FRET effect. Caspase-3 (CPP32) is an important cellular protease activated during programmed cell death. An 18 amino acid peptide contg. CPP32 recognition sequence, DEVD, was used to link GFP and BFP together. CPP32 activation can be monitored by FRET assay during the apoptosis process.

ANSWER 17 OF 18 HCAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER:

1998:94205 HCAPLUS

DOCUMENT NUMBER:

128:226014

TITLE:

Prevention of hypoxic liver cell necrosis by in vivo

human bcl-2 gene transfection

AUTHOR(S):

Yamabe, Kazuo; Shimizu, Shigeomi; Kamiike, Wataru; Waguri, Satoshi; Eguchi, Yutaka; Hasegawa, Jun-Ichi;

Okuno, Shin-Ichiro; Yoshioka, Yasuhiko; Ito,

Toshinori; Sawa, Yoshiki; Uchiyama, Yasuo; Tsujimoto,

Yoshihide; Matsuda, Hikaru

CORPORATE SOURCE:

The First Department of Surgery, Biomedical Research Center, Osaka University Medical School, Suita, 565,

Japan

SOURCE:

Biochemical and Biophysical Research Communications (

**1998**), 243(1), 217-223

CODEN: BBRCA9; ISSN: 0006-291X

PUBLISHER:

Academic Press

DOCUMENT TYPE:

Journal

LANGUAGE:

English

Prevention of hypoxic cell death is a key to successful liver transplantation. The authors developed a new method for preventing liver hypoxic cell death by introducing an anti-cell death gene directly into rat livers. When the human bcl-2 gene (hbcl-2) was directly transfected into rat livers together with nonhistone chromosomal protein high-mobility group 1 (HMG1) by the hemagglutinating virus of Japan (Sendai virus; HVJ) -liposome method, human Bcl-2 protein (hBcl-2) was efficiently expressed. Electron microscopy and

fluorescence microscopy revealed that hepatocytes expressing exogenous hBcl-2 were almost completely protected from hypoxic cell necrosis. The expression of the hBcl-2 also inhibited activation of caspase-3 (-like) proteases and liver dysfunction.

Thus, transfection of the hbcl-2 gene through HVJ-liposome method is useful to prevent liver cell necrosis induced by hypoxia. This finding could lead to new strategies to avoid the hypoxic cell death, the major problem in liver transplantation.

L7 ANSWER 18 OF 18 HCAPLUS COPYRIGHT 2002 ACS

ACCESSION NUMBER: 1998:52938 HCAPLUS

DOCUMENT NUMBER: 128:152960

TITLE: Dominant-negative FADD inhibits TNFR60-, Fas/Apol- and

TRAIL-R/Apo2-mediated cell death but not gene

induction

AUTHOR(S): Wajant, Harald; Johannes, Franz-Josef; Haas, Elvira;

Siemienski, Katrin; Schwenzer, Ralph; Schubert,

Gisela; Weiss, Tilo; Grell, Matthias; Scheurich, Peter

Institute of Cell Biology and Immunology, University

of Stuttgart, Stuttgart, 70569, Germany

Current Biology (1998), 8(2), 113-116

CODEN: CUBLE2; ISSN: 0960-9822

PUBLISHER: Current Biology Ltd.

DOCUMENT TYPE: Journal LANGUAGE: English

CORPORATE SOURCE:

SOURCE:

Fas/Apol and other cytotoxic receptors of the tumor necrosis factor ABreceptor (TNFR) family contain a cytoplasmic death domain (DD) that activates the apoptotic process by interacting with the DD-contg. adaptor proteins TNFR-assocd. DD protein (TRADD) and Fas-assocd. DD protein (FADD/MORT1), leading to the activation of cysteine proteases of the caspase family. Stimulation of Fas/Apol leads to the formation of a receptor-bound death-inducing signaling complex (DISC), consisting of FADD and two different forms of caspase-8. Transient expression of a dominant-neq. mutant of FADD impairs TNFR60-mediated and Fas/Apol-mediated apoptosis, but has no effect on TNF-related apoptosis-inducing ligand (TRAIL/Apo2L)-induced cell death. To study the function of FADD in DD-receptor signaling in more detail, we established HeLa cells that stably expressed a green fluorescent protein (GFP)-tagged dominant-neg. mutant of FADD, GFP-.DELTA.FADD. Interestingly, expression of this mutant inhibited cell death induced by TNFR60, Fas/Apo1 and TRAIL-R/Apo2. In addn., GFP-.DELTA.FADD did not interfere with TNF-mediated gene induction or with activation of NF-.kappa.B or Jun N-terminal kinase (JNK), demonstrating that FADD is part of the TNFR60-initiated apoptotic pathway but does not play a role in TNFR60-mediated gene induction. Fas/Apol-mediated activation of JNK was unaffected by the expression of GFP-.DELTA.FADD, suggesting that in Fas/Apol signaling the apoptotic pathway and the activation of JNK diverge at a level proximal to the receptor, upstream of or parallel to FADD.